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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2009; month=12; day=9; hr=8; min=45; sec=48; ms=312; ]

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Reviewer Comments:

<210> 2

<211> 205

<212> PRT

<213> artifical sequence

<400> 2

The above <213> response: 1) contains a misspelling: please change "artifical" to "Artificial". Per 1.823 of the Sequence Rules, the only valid <213> responses are: the Genus species of the organism, "Artificial Sequence", or "Unknown". 2) "Artificial Sequence" and "Unknown" require an explanation in the <220>-<223> section; please clearly indicate the source of the genetic material. If completely synthesized, please state so. Same type of errors in Sequences 3-7.

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Application No: 10526682 Version No: 2.0

Input Set:

Output Set:

Started: 2009-11-20 14:20:05.989  
Finished: 2009-11-20 14:20:06.840  
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 851 ms  
Total Warnings: 7  
Total Errors: 0  
No. of SeqIDs Defined: 7  
Actual SeqID Count: 7

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
W 402	Undefined organism found in <213> in SEQ ID (6)
W 402	Undefined organism found in <213> in SEQ ID (7)

# SEQUENCE LISTING

<110> Shi, Bingxing  
Wu, Zuze  
Yu, Aiping  
Dong, Chunna

<120> FUSED PROTEIN WITH THE FUNCTION OF BOTH HEMOLYSIS AND  
ANTICOAGULATION AND USE OF IT

<130> 0440/74021

<140> 10526682

<141> 2009-11-20

<160> 7

<170> PatentIn version 3.5

<210> 1

<211> 47

<212> DNA

<213> human

<400> 1

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47

<210> 2

<211> 205

<212> PRT

<213> artifical sequence

<400> 2

Ser Ser Ser Phe Asp Lys Gly Lys Tyr Lys Lys Gly Asp Asp Ala Ser  
1 5 10 15

Tyr Phe Glu Pro Thr Gly Pro Tyr Leu Met Val Asn Val Thr Gly Val  
20 25 30

Asp Gly Lys Gly Asn Glu Leu Leu Ser Pro His Tyr Val Glu Phe Pro  
35 40 45

Ile Lys Pro Gly Thr Thr Leu Thr Lys Glu Lys Ile Glu Tyr Tyr Val  
50 55 60

Glu Trp Ala Leu Asp Ala Thr Ala Tyr Lys Glu Phe Arg Val Val Glu  
65 70 75 80

Leu Asp Pro Ser Ala Lys Ile Glu Val Thr Tyr Tyr Asp Lys Asn Lys

85

90

95

Lys Lys Glu Glu Ser Phe Pro Ile Thr Glu Lys Gly Phe Val Val Pro  
100 105 110

Asp Leu Ser Glu His Ile Lys Asn Pro Gly Phe Asn Leu Ile Thr Lys  
115 120 125

Val Ile Ile Glu Lys Lys Gly Ser Ile Glu Gly Arg Ile Thr Tyr Thr  
130 135 140

Asp Cys Thr Glu Ser Gly Gln Asp Leu Cys Leu Cys Glu Gly Ser Asn  
145 150 155 160

Val Cys Gly Lys Gly Asn Lys Cys Ile Leu Gly Ser Asn Gly Glu Glu  
165 170 175

Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Gln Gln Ser His Asn  
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Asp Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu Gln  
195 200 205

<210> 3  
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<400> 3

Ile Glu Gly Arg  
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<210> 4  
<211> 4  
<212> PRT  
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<400> 4

Leu Gly Pro Arg  
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<210> 5  
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<212> PRT  
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<400> 5

Gly Ser Ile Glu Gly Arg  
1 5

<210> 6

<211> 6

<212> PRT

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Pro Arg Ile Glu Gly Arg  
1 5

<210> 7

<211> 6

<212> PRT

<213> artificalsequence

<400> 7

Gly Ser Leu Gly Pro Arg  
1 5